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SEQUENCE LISTING

<110> E.I. du Pont de Nemours and Company
Hallahan, David L.

<120> CIS-PRENYLTRANSFERASES FROM THE RUBBER-PRODUCING PLANTS RUSSIAN DANDELION
(TARAXACUM KOK-SAGHYZ) AND SUNFLOWER (HELIANTHUS ANNUS)

<130> CL2039

<160> 45

<170> PatentIn version 3.1

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<212> DNA

<213> Taraxacum kok-saghyz

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35 40 45

Ile Pro Glu His Ile Ala Phe Ile Leu Asp Gly Asn Arg Arg Phe Ala
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Arg Lys Trp Asn Leu Thr Glu Gly Ala Gly His Lys Thr Gly Phe Leu
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Ala Leu Met Ser Val Leu Lys Tyr Cys Tyr Glu Ile Gly Val Lys Tyr
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Val Thr Ile Tyr Ala Phe Ser Leu Asp Asn Phe Asn Arg Arg Pro Asp
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Glu Val Gln Tyr Val Met Asp Leu Met Gln Asp Lys Ile Glu Gly Phe
115 120 125

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Ile Gly Asp Leu Asp Arg Leu Tyr Glu Pro Val Arg Ile Ala Ala Glu
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Lys Ala Met Glu Ala Thr Ala Lys Asn Ser Thr Thr Tyr Leu Leu Val
165 170 175

Cys Val Ala Tyr Thr Ser Ser His Glu Ile Pro Arg Ala Ile His Glu
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Ala Cys Glu Glu Ser Ile Arg Val Met Asn Gly Asn Gly Phe Phe Asn
 195 200 205

Gly Ser Gly Tyr Thr Asn Val Asn His Gly Ser Gln Ala Val Ile Lys
 210 215 220

Val Val Asp Leu Asp Lys His Met Tyr Met Gly Val Ala Pro Asp Pro
 225 230 235 240

Asp Ile Leu Val Arg Ser Ser Gly Glu Thr Arg Leu Ser Asn Phe Leu
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Leu Trp Gln Thr Thr Asn Cys Leu Leu Tyr Ser Pro Lys Ala Leu Trp
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 35 40 45

Thr His Thr Tyr Leu Leu Val Cys Val Ala Tyr Thr Ser Ser His Glu
 50 55 60

Ile Pro Arg Ala Val Tyr Glu Ser Cys Glu Glu Lys Ser Gly Gly Thr
 65 70 75 80

Gly Val Met Ile Asn Gly Asn Gly Ser Val Asn Gly Asp Tyr Ser Glu
 85 90 95

Glu Lys Ser Gly Gly Thr Gly Val Met Val Asn Gly Asn Gly Ser Val
 100 105 110

Asn Gly Asp Tyr Ser Asn Gly Asp His Glu Glu Gly Val Lys Val Val
 115 120 125

Asp Ile Asp Lys His Met Tyr Met Ala Val Ala Pro Asp Pro Asp Ile
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Gln Thr Thr Asn Cys Val Leu Tyr
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 35 40 45

Val Ser Ile Tyr Ala Phe Ser Thr Glu Asn Trp Thr Arg Pro Lys Glu
 50 55 60
 Glu Val Asp Phe Leu Met Glu Met Tyr Glu Asp Leu Leu Arg Thr Asp
 65 70 75 80
 Ala Glu Glu Leu Leu Ser Leu Gly Cys Arg Val Ser Ile Met Gly Lys
 85 90 95
 Lys Thr Asn Leu Pro Lys Ser Leu Gln Lys Leu Cys Ile Glu Ile Glu
 100 105 110
 Glu Lys Ser Arg Ala Asn Ser Gly Thr His Val Asn Tyr Ala Leu Asn
 115 120 125
 Tyr Ser Gly Lys Tyr Asp Ile Ile Glu Ala Cys Lys Ser Val Ala Thr
 130 135 140
 Lys Val Lys Asp Gly Val Ile Ile Pro Lys Gln Ile Asp Glu Lys Tyr
 145 150 155 160
 Phe Lys Gln Glu Leu Gly Thr Lys Met Ile Asp Phe Pro Tyr Pro Asp
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 Leu Val Ile Arg Thr Ser Gly Glu Ile Arg Leu Ser Asn Phe Met Leu
 180 185 190
 Trp Gln Met Ala Tyr Ser Glu Leu Tyr Phe Thr Asp Lys Tyr Phe Pro
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 Val Arg Lys Cys
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<213> Hevea brasiliensis

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 35 40 45

Lys His Lys Leu Pro Glu Gly Gly Gly His Lys Ala Gly Phe Leu Ala
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Leu Leu Asn Val Leu Thr Tyr Cys Tyr Glu Leu Gly Val Lys Tyr Ala
 65 70 75 80

Thr Ile Tyr Ala Phe Ser Ile Asp Asn Phe Arg Arg Lys Pro His Glu
 85 90 95

Val Gln Tyr Val Met Asp Leu Met Leu Glu Lys Ile Glu Gly Met Ile
 100 105 110

Met Glu Glu Ser Ile Ile Asn Ala Tyr Asp Ile Cys Val Arg Phe Val
 115 120 125

Gly Asn Leu Lys Leu Leu Ser Glu Pro Val Lys Thr Ala Ala Asp Lys
 130 135 140

Ile Met Arg Ala Thr Ala Asn Asn Ser Lys Cys Val Leu Leu Ile Ala
 145 150 155 160

Val Cys Tyr Thr Ser Thr Asp Glu Ile Val His Ala Val Glu Glu Ser
 165 170 175

Ser Glu Leu Asn Ser Asn Glu Val Cys Asn Asn Gln Glu Leu Glu Glu
 180 185 190

Ala Asn Ala Thr Gly Ser Ser Thr Val Ile Gln Thr Glu Asn Met Glu
 195 200 205

Ser Tyr Ser Gly Ile Lys Leu Val Asp Leu Glu Lys Asn Thr Tyr Ile
 210 215 220

Asn Pro Tyr Pro Asp Val Leu Ile Arg Thr Ser Gly Glu Thr Arg Leu
 225 230 235 240

Ser Asn Tyr Leu Leu Trp Gln Thr Thr Asn Cys Ile Leu Tyr Ser Pro
 245 250 255

Tyr Ala Leu Trp Pro Glu Ile Gly Leu Arg His Val Val Trp Ser Val
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Leu Lys
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 35 40 45

Lys His Lys Leu Pro Glu Gly Gly Gly His Lys Ala Gly Phe Leu Ala
 50 55 60

Leu Leu Asn Val Leu Thr Tyr Cys Tyr Glu Leu Gly Val Lys Tyr Ala
 65 70 75 80

Thr Ile Tyr Ala Phe Ser Ile Asp Asn Phe Arg Arg Lys Pro His Glu
 85 90 95

Val Gln Tyr Val Met Asp Leu Met Leu Glu Lys Ile Glu Gly Met Ile
 100 105 110

Met Glu Glu Ser Ile Ile Asn Ala Tyr Asp Ile Cys Val Arg Phe Val
 115 120 125

Gly Asn Leu Lys Leu Leu Ser Glu Pro Val Lys Thr Ala Ala Asp Lys
 130 135 140

Ile Met Arg Ala Thr Ala Asn Asn Ser Lys Cys Val Leu Leu Ile Ala
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Val Cys Tyr Thr Ser Thr Asp Glu Ile Val His Ala Val Glu Glu Ser
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Ser Glu Leu Asn Ser Asn Glu Val Cys Asn Asn Gln Glu Leu Glu Glu
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Ala Asn Ala Thr Gly Ser Ser Thr Val Ile Gln Thr Glu Asn Met Glu
 195 200 205

Ser Tyr Ser Gly Ile Lys Leu Val Asp Leu Glu Lys Asn Thr Tyr Ile
 210 215 220

Asn Pro Tyr Pro Asp Val Leu Ile Arg Thr Ser Gly Glu Thr Arg Leu
 Page 8

225 230 235 240
 Ser Asn Tyr Leu Leu Trp Gln Thr Thr Asn Cys Ile Leu Tyr Ser Pro
 245 250 255
 Tyr Ala Leu Trp Pro Glu Ile Gly Leu Arg His Val Val Trp Ser Val
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 Leu Lys
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<213> Hevea brasiliensis

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 35 40 45
 Lys His Lys Met Lys Glu Ala Glu Gly Tyr Lys Ala Gly Tyr Leu Ala
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 Leu Leu Arg Thr Leu Thr Tyr Cys Tyr Glu Leu Gly Val Arg Tyr Val
 65 70 75 80
 Thr Ile Tyr Ala Phe Ser Ile Asp Asn Phe Arg Arg Gln Pro Arg Glu
 85 90 95
 Val Gln Cys Val Met Asn Leu Met Met Glu Lys Ile Glu Glu Ile Ile
 100 105 110
 Val Glu Glu Ser Ile Met Asn Ala Tyr Asp Val Gly Val Arg Ile Val
 115 120 125
 Gly Asn Leu Asn Leu Leu Asp Glu Pro Ile Arg Ile Ala Ala Glu Lys
 130 135 140
 Ile Met Arg Ala Thr Ala Asn Asn Ser Gly Phe Val Leu Leu Ile Ala
 145 150 155 160

Val Ala Tyr Ser Ser Thr Asp Glu Ile Gly His Ala Val Glu Glu Ser
165 170 175

Ser Lys Asp Lys Leu Asn Ser Asn Glu Val Cys Asn Asn Gly Ile Glu
180 185 190

Ala Glu Gln Glu Phe Lys Glu Ala Asn Gly Thr Gly Asn Ser Val Ile
195 200 205

Pro Val Gln Lys Thr Glu Ser Tyr Ser Gly Ile Asn Leu Ala Asp Leu
210 215 220

Glu Lys Asn Thr Tyr Val Asn Pro His Pro Asp Val Leu Ile Arg Thr
225 230 235 240

Ser Gly Leu Ser Arg Leu Ser Asn Tyr Leu Leu Trp Gln Thr Ser Asn
245 250 255

Cys Ile Leu Tyr Ser Pro Phe Ala Leu Trp Pro Glu Ile Gly Leu Arg
260 265 270

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Glu Lys His Lys Glu Tyr Leu Lys
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<212> PRT

<213> Vitis sp.

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20 25 30

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35 40 45

Thr Ala Lys Thr Asp Val Val Gly Glu Glu Glu Ala Arg Glu Val Asn
50 55 60

Glu Arg Ala Glu Glu Phe Pro Asp Gly Leu Arg Arg Glu Leu Met Pro
65 70 75 80

Glu His Val Ala Val Ile Met Asp Gly Asn Val Arg Trp Ala Gln Lys
 85 90 95
 Arg Gly Leu Pro Ala Ala Ser Gly His Gln Ala Gly Val Arg Ser Leu
 100 105 110
 Arg Glu Leu Val Glu Leu Cys Cys Lys Trp Gly Ile Lys Val Leu Ser
 115 120 125
 Val Phe Ala Phe Ser Tyr Asp Asn Trp Ser Arg Ser Glu Gly Glu Val
 130 135 140
 Gly Phe Leu Met Ser Leu Ile Glu Arg Val Val Lys Ala Glu Leu Pro
 145 150 155 160
 Ile Leu Gly Gly Lys Ala Phe Glu Cys Arg Asp Trp Gly Phe Val Lys
 165 170 175
 Ala Ser Glu Gln Leu Gln Leu Ile Ile Asp Val Glu Glu Thr Thr Lys
 180 185 190
 Glu Asn Ser Arg Leu Gln Phe Ile Val Ala Leu Ser Tyr Ser Gly Gln
 195 200 205
 Cys Asp Ile Leu Gln Ala Cys Lys Asn Ile Gly His Lys Val Lys Asp
 210 215 220
 Gly Leu Ile Glu Pro Glu Asp Ile Asn Lys Ser Leu Ile Glu Gln Glu
 225 230 235 240
 Leu Gln Thr Asn Cys Thr Glu Phe Pro Phe Pro Asp Leu Leu Ile Arg
 245 250 255
 Thr Ser Gly Glu Leu Arg Val Ser Asn Phe Met Leu Trp Gln Ile Ala
 260 265 270
 Tyr Thr Glu Leu Cys Phe Phe Ser Thr Leu Trp Pro Asp Phe Gly Lys
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<213> Oryza sativa

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 35 40 45
 Ile Thr Val Tyr Ala Phe Ser Ile Asp Asn Phe Lys Arg Asp Pro Thr
 50 55 60
 Glu Val Lys Ser Leu Met Glu Leu Met Glu Glu Lys Ile Asn Glu Leu
 65 70 75 80
 Leu Glu Asn Arg Asn Val Ile Asn Lys Val Asn Cys Lys Ile Asn Phe
 85 90 95
 Trp Gly Asn Leu Asp Met Leu Ser Lys Ser Val Arg Val Ala Ala Glu
 100 105 110
 Lys Leu Met Ala Thr Thr Ala Glu Asn Thr Gly Leu Val Phe Ser Val
 115 120 125
 Cys Met Pro Tyr Asn Ser Thr Ser Glu Ile Val Asn Ala Val Asn Lys
 130 135 140
 Val Cys Ala Glu Arg Arg Asp Ile Leu Gln Arg Glu Asp Ala Asp Ser
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 Val Ala Asn Asn Gly Val Tyr Ser Asp Ile Ser Val Ala Asp Leu Asp
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 Arg His Met Tyr Ser Ala Gly Cys Pro Asp Pro Asp Ile Val Ile Arg
 180 185 190
 Thr Ser Gly Glu Thr Arg Leu Ser Asn Phe Leu Leu Trp Gln Thr Thr
 195 200 205
 Phe Ser His Leu Gln Asn Pro Asp Pro Leu Trp Pro Glu Phe Ser Phe
 210 215 220
 Lys His Leu Val Trp Ala Ile Leu Gln Tyr Gln Arg Val His Pro Ser
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 Ile Glu Gln Ser Arg Asn Leu Ala Lys Lys Gln Leu
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<212> PRT

<213> Oryza sativa

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 35 40 45

Pro Lys His Ile Ala Phe Ile Met Asp Gly Asn Arg Arg Tyr Ala Lys
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Phe Arg Ser Ile Gln Glu Gly Ser Gly His Arg Val Gly Phe Ser Ala
 65 70 75 80

Leu Ile Ala Ser Leu Leu Tyr Cys Tyr Glu Met Gly Val Lys Tyr Ile
 85 90 95

Thr Val Tyr Ala Phe Ser Ile Asp Asn Phe Lys Arg Asp Pro Thr Glu
 100 105 110

Val Lys Ser Leu Met Glu Leu Met Glu Glu Lys Ile Asn Glu Leu Leu
 115 120 125

Glu Asn Arg Asn Val Ile Asn Lys Val Asn Cys Lys Ile Asn Phe Trp
 130 135 140

Gly Asn Leu Asp Met Leu Ser Lys Ser Val Arg Val Ala Ala Glu Lys
 145 150 155 160

Leu Met Ala Thr Thr Ala Glu Asn Thr Gly Leu Val Phe Ser Val Cys
 165 170 175

Met Pro Tyr Asn Ser Thr Ser Glu Ile Val Asn Ala Val Asn Lys Val
 180 185 190

Cys Ala Glu Arg Arg Asp Ile Leu Gln Arg Glu Asp Ala Asp Ser Val
 195 200 205

Ala Asn Asn Gly Val Tyr Ser Asp Ile Ser Val Ala Asp Leu Asp Arg
 210 215 220

His Met Tyr Ser Ala Gly Cys Pro Asp Pro Asp Ile Val Ile Arg Thr
 225 230 235 240

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Phe Glu Arg Thr Ile Asn Ser Glu Val Gln Thr Phe Lys Arg Glu Gly
165 170 175

Ile Arg Ile Ser Val Ile Gly Asp Ser Ser Arg Leu Pro Glu Ser Leu
180 185 190

Lys Arg Met Ile Ala Ser Ala Glu Glu Asp Thr Lys Gln Asn Ser Arg
195 200 205

Phe Gln Leu Ile Val Ala Val Gly Tyr Ser Gly Lys Tyr Asp Val Val
210 215 220

Gln Ala Cys Lys Ser Val Ala Lys Lys Val Lys Asp Gly His Ile His
225 230 235 240

Leu Asp Asp Ile Asn Glu Asn Ile Ile Glu Gln Glu Leu Glu Thr Asn
245 250 255

Cys Thr Glu Phe Pro Tyr Pro Asp Leu Leu Ile Arg Thr Ser Gly Glu
260 265 270

Leu Arg Val Ser Asn Phe Leu Leu Trp Gln Leu Ala Tyr Thr Glu Leu
275 280 285

Tyr Phe Asn Arg Glu Leu Trp Pro Asp Phe Gly Lys Asp Glu Phe Val
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His Ser

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<212> PRT

<213> Triticum aestivum

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 Met Arg Thr Val Arg Leu Ser Arg Ala Trp Gly Ile Arg Val Leu Thr
 65 70 75 80
 Ala Phe Gly Phe Ser Leu Glu Asn Trp Asn Arg Pro Lys Ala Glu Val
 85 90 95
 Asp Phe Leu Met Ala Leu Ile Glu Arg Phe Ile Asn Asp Asn Leu Ala
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 Glu Phe Leu Arg Glu Gly Thr Arg Leu Arg Ile Ile Gly Asp Arg Ser
 115 120 125
 Arg Leu Pro Ile Ser Val Gln Lys Thr Ala Arg Asp Ala Glu Glu Ala
 130 135 140
 Thr Arg Asn Asn Ser Gln Leu Asp Leu Val Leu Ala Ile Ser Tyr Ser
 145 150 155 160
 Gly Arg Met Asp Ile Val Gln Ala Cys Arg Asn Leu Ala Gln Lys Val
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 Asp Ala Lys Leu Leu Arg Pro Glu Asp Ile Asp Glu Ser Leu Phe Ala
 180 185 190
 Asp Glu Leu Gln Thr Ser Glu Thr Ser Cys Pro Asp Leu Leu Ile Arg
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 Thr Ser Gly Glu Leu Arg Leu Ser Asn Phe Leu Leu Trp Gln Ser Ala
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 Tyr Ser Glu Leu Phe Phe Thr Asp Thr Leu Trp Pro Asp Phe Gly Glu
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<212> PRT

<213> Dimorphotheca sinuata

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Page 17

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<301> Shimizu, N., Koyama, T. and Ogura, K.

<302> Molecular Cloning, Expression, and Purification of Undecprenyl Diphosphate Synthase: No Sequence Similarity between E- and Z-prenyl Diphosphate Synthases

<303> J. Biol. Chem.

<304> 273

<305> 31

<306> 19476-19481

<307> 1998-07-31

<308> AB004319

<309> 1997-05-29

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<213> Micrococcus luteus

<400> 18

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Arg Trp Ala Lys Gln Lys Lys Met Pro Arg Ile Lys Gly His Tyr Glu
 35 40 45

Gly Met Gln Thr Val Lys Lys Ile Thr Arg Tyr Ala Ser Asp Leu Gly
 50 55 60

Val Lys Tyr Leu Thr Leu Tyr Ala Phe Ser Thr Glu Asn Trp Ser Arg
 65 70 75 80

Pro Lys Asp Glu Val Asn Tyr Leu Met Lys Leu Pro Gly Asp Phe Leu
 85 90 95

Asn Thr Phe Leu Pro Glu Leu Ile Glu Lys Asn Val Lys Val Glu Thr
 100 105 110

Ile Gly Phe Ile Asp Asp Leu Pro Asp His Thr Lys Lys Ala Val Leu
 115 120 125

Glu Ala Lys Glu Lys Thr Lys His Asn Thr Gly Leu Thr Leu Val Phe
 130 135 140

Ala Leu Asn Tyr Gly Gly Arg Lys Glu Ile Ile Ser Ala Val Gln Leu
 145 150 155 160

Ile Ala Glu Arg Tyr Lys Ser Gly Glu Ile Ser Leu Asp Glu Ile Ser
 165 170 175

Glu Thr His Phe Asn Glu Tyr Leu Phe Thr Ala Asn Met Pro Asp Pro
 180 185 190

Glu Leu Leu Ile Arg Thr Ser Gly Glu Glu Arg Leu Ser Asn Phe Leu
 195 200 205

Ile Trp Gln Cys Ser Tyr Ser Glu Phe Val Phe Ile Asp Glu Phe Trp
 210 215 220

Pro Asp Phe Asn Glu Glu Ser Leu Ala Gln Cys Ile Ser Ile Tyr Gln
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Asn Arg His Arg Arg Phe Gly Gly Leu
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gggaacagga gattcgttag aaagaaagag atggacgtaa aggagggcca cgaggcagga    180
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aagggcgcgc ctatagacga aagcacgtta gaatcgcatc tctacacggc ggggggtaccc    600
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cctatacggg taggcatggat tttattaaaa ttttcgtttc acaaatcctt tttaaacaaa    780
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20           25           30

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Arg His Val Gly Phe Ile Met Asp Gly Asn Arg Arg Phe Ala Arg Lys
35           40           45

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Lys Glu Met Asp Val Lys Glu Gly His Glu Ala Gly Phe Val Ser Met
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Ser Arg Ile Leu Glu Leu Cys Tyr Glu Ala Gly Val Asp Thr Ala Thr
 65 70 75 80
 Val Phe Ala Phe Ser Ile Glu Asn Phe Lys Arg Ser Ser Arg Glu Val
 85 90 95
 Glu Ser Leu Met Thr Leu Ala Arg Glu Arg Ile Arg Gln Ile Thr Glu
 100 105 110
 Arg Gly Glu Leu Ala Cys Lys Tyr Gly Val Arg Ile Lys Ile Ile Gly
 115 120 125
 Asp Leu Ser Leu Leu Asp Lys Ser Leu Leu Glu Asp Val Arg Val Ala
 130 135 140
 Val Glu Thr Thr Lys Asn Asn Lys Arg Ala Thr Leu Asn Ile Cys Phe
 145 150 155 160
 Pro Tyr Thr Gly Arg Glu Glu Ile Leu His Ala Met Lys Glu Thr Ile
 165 170 175
 Val Gln His Lys Lys Gly Ala Ala Ile Asp Glu Ser Thr Leu Glu Ser
 180 185 190
 His Leu Tyr Thr Ala Gly Val Pro Pro Leu Asp Leu Leu Ile Arg Thr
 195 200 205
 Ser Gly Val Ser Arg Leu Ser Asp Phe Leu Ile Trp Gln Ala Ser Ser
 210 215 220
 Lys Gly Val Arg Ile Glu Leu Leu Asp Cys Leu Trp Pro Glu Phe Gly
 225 230 235 240
 Pro Ile Arg Met Ala Trp Ile Leu Leu Lys Phe Ser Phe His Lys Ser
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 Phe Leu Asn Lys Glu Tyr Arg Leu Glu Glu Gly Asp Tyr Asp Glu Glu
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ttaagggtag ggccagtgcc tgaacatgtc tcctttatca tggatggtaa ccggagatat      240
gccaagtcaa gaaggctacc agtaaaaaaa ggccatgaag ctggtggggt aacggtacta      300
acactactgt atatctgcaa aagattgggt gtaaaatgtg tttccgccta tgcattttct      360
attgaaaatt ttaatagacc aaaagaagaa gtagatacgc taatgaattt gtttacggta      420
aagcttgatg aattcgcaaa aagagccaag gactataagg atcccttata cggatctaaa      480
ataagaatag taggtgatca atctttacta tctccagaaa tgagaaaaaa aattaaaaaa      540
gtggaagaaa tcacacagga tggagacgat ttcactttat ttatatgttt tccttacct      600
tcaagaaatg atatgttaca tactattcgt gattcagttg aagaccattt ggaaaaataaa      660
tcaccaagga ttaatataag aaaatttact aataaaatgt acatggggtt ccattccaat      720
aaatgtgaat tattaatcag aacaagtggg cataggaggc tctcagacta tatgctatgg      780
caagtacatg aaaatgccac cattgaattt agtgatacgt tgtggccaaa ttttagcttc      840
tttgctatgt acctgatgat tctcaaatgg tccttctttt ccaccattca aaaatataat      900
gagaagaatc actcattgtt tgaaaaaata catgaaagcg ttccttcaat atttaaaaaa      960
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20          25          30
Phe Gln Arg Val Phe Ala Trp Val Met Ser Leu Ser Leu Phe Ser Trp
35          40          45
Phe Tyr Val Asn Leu Gln Asn Ile Leu Ile Lys Ala Leu Arg Val Gly
50          55          60
Pro Val Pro Glu His Val Ser Phe Ile Met Asp Gly Asn Arg Arg Tyr
65          70          75          80

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Ala Lys Ser Arg Arg Leu Pro Val Lys Lys Gly His Glu Ala Gly Gly
85 90 95

Leu Thr Leu Leu Thr Leu Leu Tyr Ile Cys Lys Arg Leu Gly Val Lys
100 105 110

Cys Val Ser Ala Tyr Ala Phe Ser Ile Glu Asn Phe Asn Arg Pro Lys
115 120 125

Glu Glu Val Asp Thr Leu Met Asn Leu Phe Thr Val Lys Leu Asp Glu
130 135 140

Phe Ala Lys Arg Ala Lys Asp Tyr Lys Asp Pro Leu Tyr Gly Ser Lys
145 150 155 160

Ile Arg Ile Val Gly Asp Gln Ser Leu Leu Ser Pro Glu Met Arg Lys
165 170 175

Lys Ile Lys Lys Val Glu Glu Ile Thr Gln Asp Gly Asp Asp Phe Thr
180 185 190

Leu Phe Ile Cys Phe Pro Tyr Thr Ser Arg Asn Asp Met Leu His Thr
195 200 205

Ile Arg Asp Ser Val Glu Asp His Leu Glu Asn Lys Ser Pro Arg Ile
210 215 220

Asn Ile Arg Lys Phe Thr Asn Lys Met Tyr Met Gly Phe His Ser Asn
225 230 235 240

Lys Cys Glu Leu Leu Ile Arg Thr Ser Gly His Arg Arg Leu Ser Asp
245 250 255

Tyr Met Leu Trp Gln Val His Glu Asn Ala Thr Ile Glu Phe Ser Asp
260 265 270

Thr Leu Trp Pro Asn Phe Ser Phe Phe Ala Met Tyr Leu Met Ile Leu
275 280 285

Lys Trp Ser Phe Phe Ser Thr Ile Gln Lys Tyr Asn Glu Lys Asn His
290 295 300

Ser Leu Phe Glu Lys Ile His Glu Ser Val Pro Ser Ile Phe Lys Lys
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Lys Lys Thr Ala Met Ser Leu Tyr Asn Phe Pro Asn Pro Pro Ile Ser
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Val ser Val Thr Gly Asp Glu
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Pro Ile Pro Thr Asn Ile Ser Phe Ile Met Asp Gly Asn Arg Arg Phe
 35 40 45

Ala Lys Lys His Asn Leu Ile Gly Leu Asp Ala Gly His Arg Ala Gly
 50 55 60

Phe Ile Ser Val Lys Tyr Ile Leu Gln Tyr Cys Lys Glu Ile Gly Val
 65 70 75 80

Pro Tyr Val Thr Leu His Ala Phe Gly Met Asp Asn Phe Lys Arg Gly
 85 90 95

Pro Glu Glu Val Lys Cys Val Met Asp Leu Met Leu Glu Lys Val Glu
 100 105 110

Leu Ala Ile Asp Gln Ala Val Ser Gly Asn Met Asn Gly Val Arg Ile
 115 120 125

Ile Phe Ala Gly Asp Leu Asp Ser Leu Asn Glu His Phe Arg Ala Ala
 130 135 140

Thr Lys Lys Leu Met Glu Leu Thr Glu Glu Asn Arg Asp Leu Ile Val
 145 150 155 160

Val Val Cys Val Ala Tyr Ser Thr Ser Leu Glu Ile Val His Ala Val
 165 170 175

Arg Lys Ser Cys Val Arg Lys Cys Thr Asn Gly Asp Asp Leu Val Leu
 180 185 190

Leu Glu Leu Ser Asp Val Glu Glu Cys Met Tyr Thr Ser Ile Val Pro
 195 200 205

Val Pro Asp Leu Val Ile Arg Thr Gly Gly Gly Asp Arg Leu Ser Asn
 210 215 220

Phe Met Thr Trp Gln Thr Ser Arg Ser Leu Leu His Arg Thr Glu Ala
 225 230 235 240

Leu Trp Pro Glu Leu Gly Leu Trp His Leu Val Trp Ala Ile Leu Lys
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Phe Gln Arg Met Gln Asp Tyr Leu Thr Lys Lys Lys Lys Leu Asp
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Gln Leu Leu Glu Gln Ile Tyr Gly Phe Ser Arg Arg Ser Leu Phe Arg
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Val Ile Ser Met Gly Pro Ile Pro Cys His Ile Ala Phe Ile Met Asp
 35 40 45

Gly Asn Arg Arg Tyr Ala Lys Lys Cys Gly Leu Leu Asp Gly Ser Gly
 50 55 60

His Lys Ala Gly Phe Ser Ala Leu Met Ser Met Leu Gln Tyr Cys Tyr
 65 70 75 80

Glu Leu Gly Ile Lys Tyr Val Thr Ile Tyr Ala Phe Ser Ile Asp Asn
 85 90 95

Phe Arg Arg Lys Pro Glu Glu Val Glu Ser Val Met Asp Leu Met Leu
 100 105 110

Glu Lys Ile Lys Ser Leu Leu Glu Lys Glu Ser Ile Val His Gln Tyr
 115 120 125

Gly Ile Arg Val Tyr Phe Ile Gly Asn Leu Ala Leu Leu Asn Asp Gln
 130 135 140

Val Arg Ala Ala Ala Glu Lys Val Met Lys Ala Thr Ala Lys Asn Ser
 145 150 155 160

Arg Val Val Leu Leu Ile Cys Ile Ala Tyr Asn Ser Thr Asp Glu Ile
 165 170 175

Val Gln Ala Val Lys Lys Ser Cys Ile Asn Lys Ser Asp Asn Ile Glu
 180 185 190

Ala Ser Asn Tyr Lys His Glu Asp Ser Asp Ser Asp Ile Glu Gly Thr
195 200 205

Asp Met Glu Asn Gln Glu Lys Lys Ile Gln Leu Val Asp Ile Glu Glu
210 215 220

Asn Met Gln Met Ser Val Ala Pro Asn Pro Asp Ile Leu Ile Arg Ser
225 230 235 240

Ser Gly Glu Thr Arg Leu Ser Asn Phe Leu Leu Trp Gln Thr Gly Asn
245 250 255

Thr Gln Leu Cys Ser Pro Ala Ala Leu Trp Pro Glu Ile Gly Leu Arg
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His Leu Leu Trp Ala Ile Leu Asn Phe Gln Arg Asn His Ser Tyr Leu
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Glu Lys Arg Lys Lys Gln Leu
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Val Phe Leu Leu Lys Leu Ile Gly Leu Ile Lys Ile Lys Ala Ala Arg
35 40 45

Asp Asn Glu Lys Arg Asp Glu Gly Thr Tyr Val Val Arg Glu Asp Gly
50 55 60

Leu Gln Arg Glu Leu Met Pro Arg His Val Ala Phe Ile Leu Asp Gly
65 70 75 80

Asn Arg Arg Trp Ala Lys Arg Ala Gly Leu Thr Thr Ser Gln Gly His
85 90 95

Glu Ala Gly Ala Lys Arg Leu Ile Asp Ile Ala Glu Leu Cys Phe Glu
100 105 110

Leu Gly Val His Thr Val Ser Ala Phe Ala Phe Ser Thr Glu Asn Trp
 115 120 125

Gly Arg Asp Lys Ile Glu Ile Asp Asn Leu Met Ser Leu Ile Gln His
 130 135 140

Tyr Arg Asn Lys Ser Asn Ile Lys Phe Phe His Arg Ser Glu Val Arg
 145 150 155 160

Val Ser Val Ile Gly Asn Lys Thr Lys Ile Pro Glu Ser Leu Leu Lys
 165 170 175

Glu Ile His Glu Ile Glu Glu Ala Thr Lys Gly Tyr Lys Asn Lys His
 180 185 190

Leu Ile Met Ala Val Asp Tyr Ser Gly Lys Phe Asp Ile Met His Ala
 195 200 205

Cys Lys Ser Leu Val Lys Lys Ser Glu Lys Gly Leu Ile Arg Glu Glu
 210 215 220

Asp Val Asp Glu Ala Leu Ile Glu Arg Glu Leu Leu Thr Asn Cys Ser
 225 230 235 240

Asp Phe Pro Ser Pro Asp Leu Met Ile Arg Thr Ser Gly Glu Gln Arg
 245 250 255

Ile Ser Asn Phe Phe Leu Trp Gln Leu Ala Tyr Ser Glu Leu Phe Phe
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Ser Pro Val Phe Trp Pro Asp Phe Asp Lys Asp Lys Leu Leu Glu Ala
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 20 25 30

Gly Thr Ser Glu Gly His Glu Ala Gly Ala Arg Arg Leu Met Glu Asn
 Page 27

35

40

45

Ala Lys Asp Cys Phe Ala Met Gly Thr Asn Thr Ile Ser Leu Phe Ala
 50 55 60

Phe Ser Thr Glu Asn Trp Glu Arg Pro Glu Asp Glu Val Lys Cys Leu
 65 70 75 80

Met Ala Leu Phe Glu Lys Tyr Leu Ala Ser Asp Met Pro Tyr Leu Arg
 85 90 95

Ser Asp Lys Ile Lys Ile Ser Val Ile Gly Asn Arg Thr Lys Leu Pro
 100 105 110

Glu Ser Leu Leu Gly Leu Ile Glu Glu Val Glu Glu Ala Thr Lys Ser
 115 120 125

Tyr Glu Gly Lys Asn Leu Ile Ile Ala Ile Asp Tyr Ser Gly Arg Tyr
 130 135 140

Asp Ile Leu Gln Ala Cys Lys Ser Leu Ala Asn Lys Val Lys Asp Gly
 145 150 155 160

Leu Ile Gln Val Glu Asp Ile Asn Glu Lys Ala Met Glu Lys Glu Leu
 165 170 175

Leu Thr Lys Cys Ser Glu Phe Pro Asn Pro Asp Leu Leu Ile Arg Thr
 180 185 190

Ser Gly Glu Gln Arg Ile Ser Asn Phe Phe Leu Trp Gln Ser Ala Tyr
 195 200 205

Thr Glu Leu Tyr Phe Pro Thr Val Leu Trp Pro Asp Phe Gly Glu Ala
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Gly Arg Arg Val

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Lys Xaa Xaa Xaa Gly
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(2):483-492)

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 1 5 10 15

Val Xaa Xaa Leu Met Xaa Leu
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(2):483-492)

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<223> xaa = D or E

<400> 33

Xaa	Leu	Xaa	Ile	Arg	Thr	Xaa	Gly	Glu	Xaa	Arg	Xaa	Ser	Asn	Phe	Xaa
1				5				10						15	

Xaa	Trp	Gln	Xaa	Xaa	Tyr	Xaa	Glu	Xaa	Xaa	Phe	Xaa	Xaa	Xaa	Xaa	Trp
-----	-----	-----	-----	-----	-----	-----	-----	-----	-----	-----	-----	-----	-----	-----	-----

20

25

30

Pro Xaa Phe
35

<210> 34

<211> 35

<212> PRT

<213> Artificial Sequence

<220>

<223> Consensus Sequence

<400> 34

Asp Ile Leu Val Arg Ser Ser Gly Glu Thr Arg Leu Ser Asn Phe Leu
1 5 10 15Leu Trp Gln Thr Thr Asn Cys Val Leu Tyr Ser Pro Lys Ala Leu Trp
20 25 30Pro Glu Met
35

<210> 35

<211> 10

<212> PRT

<213> Artificial Sequence

<220>

<223> Artificial and non-naturally occurring peptide

<400> 35

Glu Leu Val Ile Ser Leu Ile Val Glu Ser
1 5 10

<210> 36

<211> 21

<212> DNA

<213> Artificial Sequence

<220>

<223> Primer NKH46

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ttcgccggag ctccttacta a

21

<210> 37

<211> 23

<212> DNA

<213> Artificial Sequence

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<223> Primer NKH45

<400> 37
cgttcatgac ccgtatgctt tct

23

<210> 38

<211> 11

<212> PRT

<213> Artificial Sequence

<220>

<223> Consensus Sequence

<220>

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Ala Phe Ile Xaa Asp Gly Asn Arg Arg Phe Ala
1 5 10

<210> 39

<211> 10

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<220>

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<223> Xaa = T or S

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<222> (4)..(4)

<223> Xaa = any amino acid

<220>

<221> MISC_FEATURE

<222> (5)..(5)

<223> Xaa = any amino acid

<220>

<221> MISC_FEATURE

<222> (6)..(6)

<223> Xaa = D or E

<220>

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<222> (8)..(8)

<223> Xaa = any amino acid

<220>

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<222> (9)..(9)

<223> Xaa = any amino acid

<400> 39

Tyr Xaa Ser Xaa Xaa Xaa Ile Xaa Xaa Ala
1 5 10

<210> 40

<211> 22

<212> PRT

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<220>

<223> Consensus Sequence

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<223> Xaa = any amino acid

<220>

<221> MISC_FEATURE

<222> (5)..(5)

<223> Xaa = I or V

<220>

<221> MISC_FEATURE

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<223> Xaa = I or V

<220>

<221> MISC_FEATURE

<222> (9)..(9)

<223> Xaa = S or T

<220>

<221> MISC_FEATURE

<222> (12)..(12)

<223> Xaa = E or L

<220>

<221> MISC_FEATURE

<222> (13)..(13)

<223> Xaa = S or T

<220>

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<222> (18)..(18)

<223> Xaa = any amino acid

<400> 40

Pro Xaa Pro Asp Xaa Leu Xaa Arg Xaa Ser Gly Xaa Xaa Arg Leu Ser
1 5 10 15

Asn Xaa Leu Leu Trp Gln
20

<210> 41

<211> 22

<212> DNA

<213> Artificial Sequence

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<223> Primer Dan5

<400> 41
ctcgacaatt tcaatcgacg cc

22

<210> 42

<211> 22

<212> DNA

<213> Artificial Sequence

<220>

<223> Primer Dan6

<400> 42
gaaggaagtt gctcagcctt gt

22

<210> 43

<211> 21

<212> DNA

<213> Artificial Sequence

<220>

<223> Primer DegHpts

<400> 43
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21

<210> 44

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<212> DNA

<213> Artificial Sequence

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<223> n = Inosine

<220>

<221> misc_feature

<222> (8)..(8)

<223> n = Inosine

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tgccananha dgwarttrct

20

<210> 45

<211> 208

<212> PRT

<213> Taraxacum kok-saghyz

<400> 45

Leu Val Phe Ile Leu Asp Gly Asn Arg Arg Phe Ala Arg Lys Trp Asn
1 5 10 15

Leu Thr Glu Gly Ala Gly His Lys Thr Gly Phe Leu Ala Leu Met Ser
20 25 30

Val Leu Lys Tyr Cys Tyr Glu Ile Gly Val Lys Tyr Val Thr Ile Tyr
35 40 45

Ala Phe Ser Leu Asp Asn Phe Asn Arg Arg Pro Asp Glu Val Gln Tyr
50 55 60

Val Met Asp Leu Met Gln Asp Lys Ile Glu Gly Phe Leu Lys Glu Val
65 70 75 80

Ser Ile Ile Asn Gln Tyr Gly Val Arg Val Leu Phe Ile Gly Asp Leu
85 90 95

Asp Arg Leu Tyr Glu Pro Val Arg Ile Ala Ala Glu Lys Ala Met Glu
100 105 110

Ala Thr Ala Lys Asn Ser Thr Thr Tyr Leu Leu Val Cys Val Ala Tyr
115 120 125

Thr Ser Ser His Glu Ile Pro Arg Ala Ile His Glu Ala Cys Glu Glu
130 135 140

Ser Ile Arg Val Met Asn Gly Asn Gly Phe Phe Asn Gly Ser Gly Tyr
145 150 155 160

Thr Asn Val Asn His Gly Ser Gln Ala Val Ile Lys Val Val Asp Leu
165 170 175

Asp Lys His Met Tyr Met Gly Val Ala Pro Asp Pro Asp Ile Leu Val
180 185 190

Arg Ser Ser Gly Glu Thr Arg Leu Ser Asn Phe Leu Leu Trp His Lys
195 200 205